

Nam	Genotype/Features	Reference or Source
<i>P. aeruginosa</i> strains		
PAK	wild-type PAK strain	(1)
AJDP730	PAK $\Delta ctpA$	(2)
AJDP1091	PAK $\Delta lbcA::aacC1$	(2)
AJDP1140	PAK $ctpA-S302A$	(2)
<i>E. coli</i> strain		
BL21(DE3)	F- $ompT gal [dcm] [lon] hsdS_B (r_B^- m_B^-)$; <i>E. coli</i> B strain) λ DE3	(3)
Plasmids		
pET15b	Amp ^r , pMB1 <i>ori</i> , <i>T7p</i> expression vector	Novagen
pET24b	Kan ^r , pMB1 <i>ori</i> , <i>T7p</i> expression vector	Novagen
pHERD26T	Tet ^r , pMB1 <i>ori</i> , <i>araBp</i> expression vector	(4)
pQE-30	Amp ^r , Col E1 <i>ori</i> , <i>T5p</i> expression vector	Qiagen
pAJD2948	<i>T5p-his₆-PA1198</i> in pQE-30	(5)
pAJD3037	<i>araBp-ctpA</i> in pHERD26T	This study
pAJD3038	<i>araBp-ctpAΔC6</i> in pHERD26T	This study
pAJD3039	<i>araBp-ctpA-L426K L430K</i> in pHERD26T	This study
pAJD3045	<i>araBp-ctpA-S302A</i> in pHERD26T	This study
pAJD3063	<i>araBp-ctpA-L426A L430A</i> in pHERD26T	This study
pAJD3109	<i>araBp-lbcA</i> in pHERD26T	This study
pAJD3110	<i>araBp-lbcAΔN84</i> in pHERD26T	This study
pAJD3111	<i>araBp-lbcAΔN165</i> in pHERD26T	This study
pET15b_CtpAΔN37	<i>T7p-his₆-ctpAΔN37</i> in pET15b	This study
pET15b_CtpAΔN37ΔC6	<i>T7p-his₆-ctpAΔN37ΔC6</i> in pET15b	This study
pET15b_CtpAΔN37_S302A	<i>T7p-his₆-ctpAΔN37_S302A</i> in pET15b	This study

pET15b_CtpAΔN37_L426K/L430K	<i>T7p-his₆-ctpAΔN37_L426K L430K</i> in pET15b	This study
pET15b_CtpAΔN37_L426A/L430A	<i>T7p-his₆-ctpAΔN37_L426A L430A</i> in pET15b	This study
pET24b_LbcAΔN48	<i>T7p-lbcAΔN48-his₆</i> in pET24b	This study
pET15b_LbcAΔN31	<i>T7p-his₆-lbcAΔN31</i> in pET15b	This study
pET15b_LbcAΔN48	<i>T7p-his₆-lbcAΔN48</i> in pET15b	This study
pET15b_LbcAΔN84	<i>T7p-his₆-lbcAΔN84</i> in pET15b	This study
pET15b_LbcAΔN165	<i>T7p-his₆-lbcAΔN165</i> in pET15b	This study
pET15b_LbcAΔN48ΔC5	<i>T7p-his₆-lbcAΔN48ΔC5</i> in pET15b	This study

REFERENCES

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4. Qiu D, Damron FH, Mima T, Schweizer HP, Yu HD. 2008. PBAD-based shuttle vectors for functional analysis of toxic and highly regulated genes in *Pseudomonas* and *Burkholderia* spp. and other bacteria. Appl Environ Microbiol 74:7422-7426.
5. Chakraborty D, Darwin AJ. 2021. Direct and indirect interactions promote complexes of the lipoprotein LbcA, the CtpA protease and its substrates, and other cell wall proteins in *Pseudomonas aeruginosa*. J Bacteriol doi:10.1128/JB.00393-21:JB0039321.